## 32-4391: Recombinant Human Poly(A) Binding Protein, Nuclear 1

Alternative Name :

Poly(A) Binding Protein Nuclear 1,Poly(A) Binding Protein 2,Polyadenylate-Binding Nuclear Protein 1,PABP-2,OPMD,PAB2,Nuclear Poly(A)-Binding Protein 1,PABII.

## Description

Source : Escherichia Coli. PABPN1 Human Recombinant produced in E.coli is a single, non-glycosylated polypeptide chain containing 211 amino acids (119-306) and having a molecular mass of 23.8 kDa . PABPN1 is fused to a 23 amino acid His-tag at $N$-terminus. PABPN1 is an highly expressed nuclear protein which has high affinity to nascent poly $(A)$ tails. PABPN1 is essential for progressive and efficient polymerization of $\operatorname{poly}(\mathrm{A})$ tails at the $3^{\prime}$ ends of eukaryotic transcripts and regulates the length of the poly $(A)$ tail to about 250 nt . In its steady-state, PABPN1 is restricted to the nucleus however there are different poly $(A)$ binding proteins which can be found in the cytoplasm. PABPN1 has a GCG trinucleotide repeat at the $5^{\prime}$ end of the coding region. Expansion of this repeat from the normal 6 copies to $8-13$ copies results in autosomal dominant oculopharyngeal muscular dystrophy (OPMD) disease. Associated pseudogenes were located on chromosomes 19 and X . Additionally, there is a read-through transcription between PABPN1 gene and the adjacent upstream BCL2L2 (BCL2-like 2) gene.

## Product Info

## Amount :

## Purification :

## Content:

## Storage condition :

Amino Acid :
$20 \mu \mathrm{~g}$
Greater than $95 \%$ as determined by SDS-PAGE.
The PABPN1 solution ( $1 \mathrm{mg} / \mathrm{ml}$ ) contains 20 mM Tris-HCl buffer ( pH 8.0 ), 0.4 M Urea and $10 \%$ glycerol.
Store at $4^{\circ} \mathrm{C}$ if entire vial will be used within 2-4 weeks. Store, frozen at $-20^{\circ} \mathrm{C}$ for longer periods of time. For long term storage it is recommended to add a carrier protein ( $0.1 \% \mathrm{HSA}$ or BSA).Avoid multiple freeze-thaw cycles.
MGSSHHHHHH SSGLVPRGSH MGSLEAIKAR VREMEEEAEK LKELQNEVEK QMNMSPPPGN AGPVIMSIEE KMEADARSIY VGNVDYGATA EELEAHFHGC GSVNRVTILC DKFSGHPKGF AYIEFSDKES VRTSLALDES LFRGRQIKVI PKRTNRPGIS TTDRGFPRAR YRARTTNYNS SRSRFYSGFN SRPRGRVYRG RARATSWYSP Y


